Predictive Ranking	Residue	Amino Acid Sequence	Secondary Structure
7	27 – 45	Tgtdmklrlpaspethldm	25 – 28 β turn; 29 – 32 α helix; 35 – 38 β turn
8 (DW5)	115 – 136	AVLDNGDPL**NNTT**PVTGASPGG	116 – 135 β turn
9	168 – 189	LWKDIFHKNNQLALTLIDT**NRS**	173 – 176 β turn; 177 – 181 α helix
1	182 – 216	TLIDT**NRSR**ACHPCSPMCKGSRCWG ESSEDCQSLT	184 – 212 β turn/loop
6	270 – 290	ALVTYNTDTFESMPNPEGRYT	273 – 286 β turn; 278 – 280 α helix
3	316 – 339	PLHNQEVTAEDGTQRAEKCSKPCA	319 – 324 α helix; 324 – 336 β turn.
10 (DW1)	376 – 395	PESFDGDPASNTAPLQPE	379 – 388 β turn
12 (DW6)	410-429	LYISAWPDSLPDLSVFQNLQ	413-421 β turn
2	485 – 503	LFRNPHQALLHTANRPEDE	497 – 500 β turn; 499 – 504 α helix
11	560 – 593	CLPCHPECQPQ**NGSV**TCFGPEADQCVACAH YKDP	561 – 572 & 589 – 593 β turn; 579 – 581 α helix
4	605– 622	KPDLSYMPIWKFPDEEGA	616 – 620 α helix
5 (DW4)	628 – 650	IN*G*THSCVDLDDKGCPAEQRASP	635 – 642 β turn; 643 – 646 α helix

Page 36, line 6:

Please replace the Sequence Listings which were filed on April 9, 2001, November 21, 2001, and April 1, 2002 with the Sequence Listing which is attached hereto.

IN THE CLAIMS:

1. (Once Amended) A composition for stimulating an immune response to HER-2 protein, wherein said composition is a chimeric peptide and comprises a HER-2 B cell epitope, a T helper (Th) epitope, and a linker joining said HER-2 B cell epitope to said Th epitope; said HER-2 B cell epitope being from 15 to 40 amino acids in length and comprising a sequence selected from the group consisting of:

TGTDMKLRLPASPETHLDM, SEQ ID NO. 1, or a functional equivalent thereof;

TLIDTNRSRACHPCSPMCKGSRCWGESSEDCQSLT, SEQ ID NO. 4, or a functional equivalent thereof;

ALVTYNTDTFESMPNPEGRYT, SEQ ID NO. 5, or a functional equivalent thereof;

PLHNQEVTAEDGTQRAEKCSKPCA, SEQ ID NO. 6, or a functional equivalent thereof;

LFRNPHQALLHTANRPEDE, SEQ ID NO. 9, or a functional equivalent thereof;